BIOTECHN... JGY
SYSTEMS
BRANCH
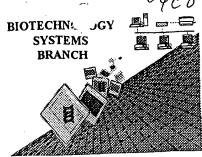
# RAW SEQUENCE LISTING
# ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:  _09/776,724_

Source:  _OIPE_

Date Processed by STIC:  _2/28/2001_

**THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.**
**PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:**
1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**
**FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.**

**FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.**
**PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)**
**PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)**

**TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:**

## Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.
Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).
Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:
## http://www.uspto.gov/web/offices/pac/checker

RAW SEQUENCE LISTING
PATENT APPLICATION:  US/09/776,724

DATE: 02/28/2001
TIME: 15:50:50

Input Set : A:\PZ011_SeqList.txt
Output Set: N:\CRF3\02282001\I776724.raw

Does Not Comply
Corrected Diskette Needed

pp 1-2,5

```
        2 <110> APPLICANT: Rosen et al.
        4 <120> TITLE OF INVENTION: 64 Human Secreted Proteins
        6 <130> FILE REFERENCE: PZ011
C-->    8 <140> CURRENT APPLICATION NUMBER: US/09/776,724
        9 <141> CURRENT FILING DATE: 2001-02-06
       11 <150> PRIOR APPLICATION NUMBER: 60/180,909
       12 <151> PRIOR FILING DATE: 2000-02-08
       14 <150> PRIOR APPLICATION NUMBER: 09/669,688
       15 <151> PRIOR FILING DATE: 2000-09-26
       17 <150> PRIOR APPLICATION NUMBER: 09/229,982
       18 <151> PRIOR FILING DATE: 1999-01-14
       20 <150> PRIOR APPLICATION NUMBER: 1998-07-15
W-->   21 <151> PRIOR FILING DATE: PCT/US98/14613
       23 <150> PRIOR APPLICATION NUMBER: 1997-07-16
W-->   24 <151> PRIOR FILING DATE: 60/052,661
W-->   26 <150> PRIOR APPLICATION NO: 1997-07-16
W-->   27 <151> PRIOR FILING DATE: 60/052,872
W-->   29 <150> PRIOR APPLICATION NO: 1997-07-16
W-->   30 <151> PRIOR FILING DATE: 60/052,871
W-->   32 <150> PRIOR APPLICATION NO: 1997-07-16
W-->   33 <151> PRIOR FILING DATE: 60/052,874
W-->   35 <150> PRIOR APPLICATION NO: 1997-07-16
W-->   36 <151> PRIOR FILING DATE: 60/052,873
W-->   38 <150> PRIOR APPLICATION NO: 1997-07-16
W-->   39 <151> PRIOR FILING DATE: 60/052,870
W-->   41 <150> PRIOR APPLICATION NO: 1997-07-16
W-->   42 <151> PRIOR FILING DATE: 60/052,875
       44 <150> PRIOR APPLICATION NUMBER: 1997-07-22
W-->   45 <151> PRIOR FILING DATE: 60/053,440
W-->   47 <150> PRIOR APPLICATION NO: 1997-07-22
W-->   48 <151> PRIOR FILING DATE: 60/053,441
W-->   50 <150> PRIOR APPLICATION NO: 1997-07-22
W-->   51 <151> PRIOR FILING DATE: 60/053,442
       53 <150> PRIOR APPLICATION NUMBER: 1997-08-18
W-->   54 <151> PRIOR FILING DATE: 60/056,359
W-->   56 <150> PRIOR APPLICATION NO: 1997-08-18
W-->   57 <151> PRIOR FILING DATE: 60/055,725
W-->   59 <150> PRIOR APPLICATION NO: 1997-08-18
W-->   60 <151> PRIOR FILING DATE: 60/055,985
W-->   62 <150> PRIOR APPLICATION NO: 1997-08-18
W-->   63 <151> PRIOR FILING DATE: 60/055,952
W-->   65 <150> PRIOR APPLICATION NO: 1997-08-18
W-->   66 <151> PRIOR FILING DATE: 60/055,989
W-->   68 <150> PRIOR APPLICATION NO: 1997-08-18
W-->   69 <151> PRIOR FILING DATE: 60/056,361
W-->   71 <150> PRIOR APPLICATION NO: 1997-08-18
W-->   72 <151> PRIOR FILING DATE: 60/055,726
```

<150> is Application No:

<151> is Application filing date:

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/776,724

DATE: 02/28/2001
TIME: 15:50:50

Input Set : A:\PZ011_SeqList.txt
Output Set: N:\CRF3\02282001\I776724.raw

W--> 74 <150> PRIOR APPLICATION NO: 1997-08-18
W--> 75 <151> PRIOR FILING DATE: 60/055,724
W--> 77 <150> PRIOR APPLICATION NO: 1997-08-18
W--> 78 <151> PRIOR FILING DATE: 60/055,946
W--> 80 <150> PRIOR APPLICATION NO: 1997-08-18
W--> 81 <151> PRIOR FILING DATE: 60/055,683

*See page 1*

```
 83 <160> NUMBER OF SEQ ID NOS: 294
 85 <170> SOFTWARE: PatentIn Ver. 2.0
 88 <210> SEQ ID NO: 1
 89 <211> LENGTH: 733
 90 <212> TYPE: DNA
 91 <213> ORGANISM: Homo sapiens
 93 <400> SEQUENCE: 1
 94 gggatccgga gcccaaatct tctgacaaaa ctcacacatg cccaccgtgc ccagcacctg     60
 95 aattcgaggg tgcaccgtca gtcttcctct tccccccaaa acccaaggac accctcatga    120
 96 tctcccggac tcctgaggtc acatgcgtgg tggtggacgt aagccacgaa gaccctgagg    180
 97 tcaagttcaa ctggtacgtg gacggcgtgg aggtgcataa tgccaagaca aagccgcggg    240
 98 aggagcagta caacagcacg taccgtgtgg tcagcgtcct caccgtcctg caccaggact    300
 99 ggctgaatgg caaggagtac aagtgcaagg tctccaacaa agccctccca acccccatcg    360
100 agaaaaccat ctccaaagcc aaagggcagc cccgagaacc acaggtgtac accctgcccc    420
101 catcccggga tgagctgacc aagaaccagg tcagcctgac ctgcctggtc aaaggcttct    480
102 atcccagcga catcgccgtg gagtgggaga gcaatgggca gccggagaac aactacaaga    540
103 ccacgcctcc cgtgctggac tccgacggct ccttcttcct ctacagcaag ctcaccgtgg    600
104 acaagagcag gtggcagcag gggaacgtct tctcatgctc cgtgatgcat gaggctctgc    660
105 acaaccacta cacgcagaag agcctctccc tgtctccggg taaatgagtg cgacggccgc    720
106 gactctagag gat                                                       733
109 <210> SEQ ID NO: 2
110 <211> LENGTH: 5
111 <212> TYPE: PRT
112 <213> ORGANISM: Homo sapiens
114 <220> FEATURE:
115 <221> NAME/KEY: Site
116 <222> LOCATION: (3)
117 <223> OTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino acids
119 <400> SEQUENCE: 2
120 Trp Ser Xaa Trp Ser
121  1               5
123 <210> SEQ ID NO: 3
124 <211> LENGTH: 86
125 <212> TYPE: DNA
126 <213> ORGANISM: Homo sapiens
128 <400> SEQUENCE: 3
129 gcgcctcgag atttccccga aatctagatt tccccgaaat gatttccccg aaatgatttc     60
130 cccgaaatat ctgccatctc aattag                                         86
133 <210> SEQ ID NO: 4
134 <211> LENGTH: 27
135 <212> TYPE: DNA
136 <213> ORGANISM: Homo sapiens
138 <400> SEQUENCE: 4
```

OK>

```
                                                                                  27

139 gcggcaagct ttttgcaaag cctaggc
142 <210> SEQ ID NO: 5
143 <211> LENGTH: 271
144 <212> TYPE: DNA
145 <213> ORGANISM: Homo sapiens                                           60
147 <400> SEQUENCE: 5
148 ctcgagattt ccccgaaatc tagatttccc cgaaatgatt tccccgaaat gatttccccg      120
149 aaatatctgc catctcaatt agtcagcaac catagtcccg cccctaactc cgcccatccc      180
150 gcccctaact ccgcccagtt ccgcccattc tccgccccat ggctgactaa ttttttttat      240
151 ttatgcagag gccgaggccg cctcggcctc tgagctattc cagaagtagt gaggaggctt      271
152 ttttggaggc ctaggctttt gcaaaaagct t
155 <210> SEQ ID NO: 6
156 <211> LENGTH: 32
157 <212> TYPE: DNA
158 <213> ORGANISM: Homo sapiens                                           32
160 <400> SEQUENCE: 6
161 gcgctcgagg gatgacagcg atagaacccc gg
164 <210> SEQ ID NO: 7
165 <211> LENGTH: 31
166 <212> TYPE: DNA
167 <213> ORGANISM: Homo sapiens                                           31
169 <400> SEQUENCE: 7
170 gcgaagcttc gcgactcccc ggatccgcct c
173 <210> SEQ ID NO: 8
174 <211> LENGTH: 12
175 <212> TYPE: DNA
176 <213> ORGANISM: Homo sapiens                                           12
178 <400> SEQUENCE: 8
179 ggggactttc cc
182 <210> SEQ ID NO: 9
183 <211> LENGTH: 73
184 <212> TYPE: DNA
185 <213> ORGANISM: Homo sapiens                                           60
187 <400> SEQUENCE: 9
188 gcggcctcga ggggactttc ccggggactt tccgggact tccatcctg                  73
189 ccatctcaat tag
192 <210> SEQ ID NO: 10
193 <211> LENGTH: 256
194 <212> TYPE: DNA
195 <213> ORGANISM: Homo sapiens                                           60
197 <400> SEQUENCE: 10
198 ctcgagggga ctttcccggg gactttccgg ggactttccg ggactttcca tctgccatct     120
199 caattagtca gcaaccatag tcccgcccct aactccgccc atcccgcccc taactccgcc     180
200 cagttccgcc cattctccgc cccatggctg actaattttt tttatttatg cagaggccga     240
201 ggccgcctcg gcctctgagc tattccagaa gtagtgagga ggctttttg gaggcctagg      256
202 cttttgcaaa aagctt
205 <210> SEQ ID NO: 11
206 <211> LENGTH: 558
207 <212> TYPE: DNA
```

2/28/01

```
208 <213> ORGANISM: Homo sapiens
210 <220> FEATURE:
211 <221> NAME/KEY: SITE
212 <222> LOCATION: (546)
213 <223> OTHER INFORMATION: n equals a,t,g, or c
215 <400> SEQUENCE: 11
216 gaattcggca cgagctgggc tgcagttggc gattccgcgc ggtgaaagca gccagtgccc     60
217 agggtctttt cctgagtgca cctgggcctg ccgcccggcg atgccatggg gtcgtgcgct    120
218 gcttttctac ttgccgcgct ctcactgctc ggtgtactgg gagggtaccc tgggaggcgt    180
219 gcctttattc ttccgaaccg ccgctcactg agacagtggc tagaagtgtc tcttggacct    240
220 gtgagttagc cttaacctgt tatgccccca gagccctcag tggagcgccc gtactttgcc    300
221 ggcatgacgt ttgatttccc ggtgataatc cgacgagttt gacagattga ggtagtgagc    360
222 aaagttgccc gtcagttggt ggccacttga cttcgtgcgg accctggcct tgctcttgga    420
223 agagatagtg ttcttagggc tggttttcact gtctcttaag actgaarggt ggarctggga    480
224 tatagatgtg ttgtttcttt tcaaatcaaa cctgcttarg tcgtcactcg aaggggggcc    540
```
OK 225 cggtanccaa ttcgccct    558

```
228 <210> SEQ ID NO: 12
229 <211> LENGTH: 715
230 <212> TYPE: DNA
231 <213> ORGANISM: Homo sapiens
233 <400> SEQUENCE: 12
234 cggatttcga gtgcttttct ccttacctcc accctccccc atgttttaat gcagccctcc     60
235 aaaaatattt actgagtgtg gactctagac cagggcctgt gctaggatac aaagatgaat    120
236 gaggcaccac ccttatcttc gagtagtata tgttttattt tattttattt ttttcccctg    180
237 ctgcctccct tgagtagtac atgttttagt aaggggaaca gacactaaag agtcctggta    240
238 atgatgagca aagtactgca tgagtaagta tctggggggc aagtgtcccc actaggactc    300
239 ctgtcagatc tggaaaaggc ctgaggaatc tgatacatga cttaatgcag cgtatacttg    360
240 cagcctggaa aactaagtaa tgacaaaata gacattcttg tcagtgtgag ccattctctg    420
241 agtccmaggg gagtacataa ttcaaaccag aattggtcat tttggagttt gcactcttag    480
242 cagtatacag tggagtgaaa tttaagaatc aatttaattt cttttcagtt tttatgtaca    540
243 taaaacctgc ttactacaag agacccagtt tattattttg tgttggttaa cattcataag    600
244 tatatttcat cataataagg ctccgtgaaa ttagtcattt tatcatttgc caataaagac    660
245 atatatctga aaataaatgt tcctgaacct gaaaaaaaaa aaaaaaaaaa ctcga         715
```

```
248 <210> SEQ ID NO: 13
249 <211> LENGTH: 838
250 <212> TYPE: DNA
251 <213> ORGANISM: Homo sapiens
253 <400> SEQUENCE: 13
254 gaattcggca cgagccaaaa caaaagaaac ctttggaggc atgtgtcaga acagagaaag     60
255 tgtcctggtt ttgcttatag aatcaaatat gttctcattc tacctactgt tttcattcta    120
256 catagtgttt tccttcttta tagtttttacg tcctcttcct aggaatgagt ctattaagaa    180
257 aataggtgtt atctttttagc tttggcattt gactttcagg ataatagagc tatctgctac    240
258 tgacagaaaa gctttgacaa gtgtttaata ctctgggatt accttcatct tactttttgca    300
259 atcattatgt gaacattgtc ttccgtccac atctayaggc tagtawgtaa caccgttgac    360
260 taaatccaaa ctttaggcta gggaaaaagg gtatactttc tgggtttcgg ttgtagatta    420
261 tgtttagatc taaycaaaac aggacagtgg tccaaacaga aaattgctat tttctgtatc    480
262 ttgtaaatct aggatttgag tttttaagat gaatttatgg ttccctttct gatatcattt    540
263 ctcatctgca gctcctaatg cctggtacct tgggtatgga gtgaggagag acaatggaca    600
264 gttttatata agaaatggaa gtaatgatac tatctttcct ggaatatttg caggccccag    660
```

```
265 aggagatgat gagcaaggac tgttggcctg tattacacac aacagggttg tagttactat      720
266 cccagcaagg aaagggtgta tctttcttct ttcatgcaaa ttatctatga tgacctaaca      780
267 gtttgattat agtgagtgga ctaaccacaa caataaaaaa aaaaaaaaaa aactcgaa        838
270 <210> SEQ ID NO: 14
271 <211> LENGTH: 513
272 <212> TYPE: DNA
273 <213> ORGANISM: Homo sapiens
275 <400> SEQUENCE: 14
276 ctgcaggaat tcggcacgag ggaacaactc catgtttttg taaaggccta gagaacatat       60
277 atccagtgcc tttccttttt gcctttgtat tcatcatttt ggcaaattac tggaagatga      120
278 cggttctggc caaaaggctg gttttgtttt tgggtcacat tttcttgctt ctctgcgtta      180
279 gaatcttgga ttagatgatg gacatggtga agatctcagc aacctcattc actagaagat      240
280 catgtggatt ggaatcatac aatggggaac aaatggaaaa gagtactttt gaaatagtgc      300
281 tggagaccac tgtgaccaca gaatgtcaag acacgtgctg ccattactgt tactatttgg      360
282 aaaatacatt cttgtaaatg caaccttagg gggtttgagg gggaagtctg ttgggaaatg      420
283 aattgcaaga aaaatattac accctgaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa      480
284 aaaaaaaaaa aaaaaaaaaa aaaaaaaact cga                                  513
287 <210> SEQ ID NO: 15
288 <211> LENGTH: 712
289 <212> TYPE: DNA
290 <213> ORGANISM: Homo sapiens
292 <220> FEATURE:
293 <221> NAME/KEY: SITE
294 <222> LOCATION: (565)
295 <223> OTHER INFORMATION: n equals a,t,g, or c
297 <400> SEQUENCE: 15
298 gaattcggca cgagctggac aggaccggag aggaccccgc gtaaccgcgg aacagacact       60
299 cccggcagcg gccgccgccg cggcactgct acgggacgag ccggagcgct tggccatggc      120
300 ggcccgatcc gcactggcgc tgctgctgct gctgccagtc ctgctcctgc cggtgcagag      180
301 ycgctcagag cccgagacca ccgcgcccac ccctacccca atcccgggtg gcaactcgtc      240
302 aktgagcagg cccctgccca gcatcgagct ccacgcctgc ggcccatacc ccaaaccagc      300
303 cctgctcatc ctgctggccc cgctggccct gtggcccatt ctcctgtagg gacgcccagc      360
304 cagccacctc taagtcgccg ctgggactgg cctgccccat tgagcaacag agacgcttga      420
305 cagccgcccg cctccattcc ttgacttcac ccagaaatgg gtccagaaaa ctgaatccca      480
306 ccagcactgg tttggagcaa ccggacaccg aggtttcacc tccagggrtt ccatggaaga      540
307 gcctcaatgg agatgccaca tcctnactga gttaaagatg ggctgaggaa cttgggtacc      600
308 cacaagtytg ccttgggrat caaaagaaaa tatttacctt tagtttggtt cattaaatgc      660
309 atgaagtcaa aatatgaaaa aaaaaaaaaa aaaaaaaaaa aaaaaactc ga               712
312 <210> SEQ ID NO: 16
313 <211> LENGTH: 652
314 <212> TYPE: DNA
315 <213> ORGANISM: Homo sapiens
317 <400> SEQUENCE: 16
318 gaattcggca cgagcaacag tggggcactc tgctcccagg caggtcccac tgggctgagc       60
319 cgcacagcct ggctttgggc ttccctgact gcaccaccca catcasctgc ctctagccct      120
320 taamatacaa aacttccccc agtcactggc cgccaggctg agttggggga tgtgttacat      180
321 ccctgggtcc actgggggc agtgttggcc atggtgttgg tgctggctct gccgagaggc      240
322 gttggagtgg ctgtgtgggg cggtgagcgc cggcccagcc tgatggaacc cactgtacca      300
323 ggcccaggcc tcagcctctg agaaggactt ccctgtgtca ctcactcata catgtcctca      360
```

2/28/01

VERIFICATION SUMMARY
PATENT APPLICATION:  US/09/776,724

DATE: 02/28/2001
TIME: 15:50:51

Input Set : A:\PZ011_SeqList.txt
Output Set: N:\CRF3\02282001\I776724.raw

```
L:8 M:270 C: Current Application Number differs, Replaced Current Application Number
L:21 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:24 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:26 M:288 W: Application Number is Repeated, <150> PRIOR APPLICATION NUMBER
L:27 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:29 M:288 W: Application Number is Repeated, <150> PRIOR APPLICATION NUMBER
L:30 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:32 M:288 W: Application Number is Repeated, <150> PRIOR APPLICATION NUMBER
L:33 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:35 M:288 W: Application Number is Repeated, <150> PRIOR APPLICATION NUMBER
L:36 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:38 M:288 W: Application Number is Repeated, <150> PRIOR APPLICATION NUMBER
L:39 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:41 M:288 W: Application Number is Repeated, <150> PRIOR APPLICATION NUMBER
L:42 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:45 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:47 M:288 W: Application Number is Repeated, <150> PRIOR APPLICATION NUMBER
L:48 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:50 M:288 W: Application Number is Repeated, <150> PRIOR APPLICATION NUMBER
L:51 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:54 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:56 M:288 W: Application Number is Repeated, <150> PRIOR APPLICATION NUMBER
L:57 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:59 M:288 W: Application Number is Repeated, <150> PRIOR APPLICATION NUMBER
L:60 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:62 M:288 W: Application Number is Repeated, <150> PRIOR APPLICATION NUMBER
L:63 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:65 M:288 W: Application Number is Repeated, <150> PRIOR APPLICATION NUMBER
L:66 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:68 M:288 W: Application Number is Repeated, <150> PRIOR APPLICATION NUMBER
L:69 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:71 M:288 W: Application Number is Repeated, <150> PRIOR APPLICATION NUMBER
L:72 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:74 M:288 W: Application Number is Repeated, <150> PRIOR APPLICATION NUMBER
L:75 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:77 M:288 W: Application Number is Repeated, <150> PRIOR APPLICATION NUMBER
L:78 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:80 M:288 W: Application Number is Repeated, <150> PRIOR APPLICATION NUMBER
L:81 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:120 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:225 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:307 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:399 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:420 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:489 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:545 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:566 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:660 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
```

VERIFICATION SUMMARY
PATENT APPLICATION:  US/09/776,724

DATE: 02/28/2001
TIME: 15:50:51

Input Set : A:\PZ011_SeqList.txt
Output Set: N:\CRF3\02282001\I776724.raw

```
L:662  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:663  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:809  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33
L:850  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34
L:851  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34
L:877  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:906  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36
L:948  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38
L:981  M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39
L:1024 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40
L:1087 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42
L:1089 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42
L:1090 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42
L:1113 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43
L:1133 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44
L:1189 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45
L:1208 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:46
L:1233 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47
L:1251 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47
L:1375 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:1423 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:53
L:1449 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:54
L:1515 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56
L:1516 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56
L:1517 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56
L:1541 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:57
L:1645 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60
L:1724 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:64
L:1802 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67
L:1812 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67
L:1813 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67
L:1861 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:68
L:1887 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:69
L:1984 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:73
L:2139 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78
L:2141 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78
L:2142 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78
L:2173 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:79
L:2251 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81
L:2271 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81
L:2379 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:87
L:4574 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=191
```

2/28/01